Amendments to the Claims

This listing of claims will replace all prior versions, and listings, of claims in the application:

1.-38. (canceled)

- 39. (previously presented) A cell that produces a glycoconjugate of interest in the absence of an exogenously supplied nucleotide triphosphate, the cell comprising heterologous genes encoding one or more sugar nucleotide regenerating enzyme and one or more glycosyltransferase.
- 40. (original) The cell of claim 39, wherein the cell is a prokaryotic cell.
- 41. (original) The cell of claim 40, wherein the prokaryotic cell is a bacterium.
- 42. (original) The cell of claim 41, wherein the bacterium is *E. coli*.
- 43. (original) The cell of claim 42, wherein the E. coli is Lac Z^- .
- 44. (original) The cell of 39, wherein the cell is a eukaryotic cell.
- 45. (original) The cell of claim 44, wherein the eukaryotic cell is a yeast.
- 46. (original) The cell of claim 39, wherein at least one of the heterologous genes is integrated into the genome of the cell.
- 47. (previously presented) The cell of claim 39, wherein the heterologous genes are encoded within one or more plasmids.
- 48. (previously presented) The cell of claim 47, wherein the heterologous genes are encoded within one plasmid.

49. (withdrawn) A method of producing a glycoconjugate of interest in the absence of an exogenously supplied nucleotide triphosphate, comprising the step of contacting a cell comprising heterologous genes encoding:

- (i). one or more encoding sugar nucleotide regenerating enzymes selected from the group consisting of GalK, GaIT, GaIU, PykF, Ndk, PpK, AcK, PoxB, Ppa, PgM, NagE, Agm1, glmU, a GaINAc kinase, a pyrophosphorylase, Ugd, NanA, Cmk, NeuA, AIg2, AIg1, SusA, ManB, ManC, a phosphomannomutase, GaIE, GMP, GMD, and GFS; and
- (ii). one or more glycosyltransferase, with a bioenergetic.

50.-51. (canceled)

(currently amended) The cell of claim 39, wherein the one or more sugar 52. nucleotide regenerating enzyme is selected from the group consisting of galactokinase, GalK, galactose-1-phosphate uridylyltransferase, GalT, glucose-1phosphate uridylyltransferase, GalU, pyruvate kinase, PykF, nucleotide diphosphate kinase, Ndk, polyphosphate kinase, PpK, acetate kinase, AcK, pyruvate oxidase, PoxB, pyrophosphatase, Ppa, phosphoglucomutase, PgM, Nacetylglucosamine permease, NagE, acetylglucosamine-phosphate mutase, Agm1, N-acetylglucosamine-1-phosphate uridyltransferase, glmU, a Nacetylglucosamine GalNAe kinase, a pyrophosphorylase, uridine 5'diphosphoglucouronic acid 6-dehydrogenase, Ugd, N-acetylneuraminate lyase, sialic acid aldolase, NanA, cytosine 5'-monophosphate kinase, Cmk, cytosine 5'monophosphate-N-acetylneuraminic acid synthetase, NeuA, α1,3mannosyltransferase, Alg2, guanosine 5'-diphosphomannose:Dol-PP-Nacetylglucosamine β-mannosyltransferase, Alg1, sucrose synthetase, SusA, ManB, mannose-1-phosphate guanyltransferase, guanisine 5'-diphosphate-mannose pyrophosphorlyase, ManC, a phosphomannomutase, uridine 5'-diphosphategalactose 4-epimease, uridine 5'-diphosphate-glucose 4-epimerase, GalE, guanosine 5'-diphosphate-mannose pyrophosphorylase, GMP, guanosine 5'diphosphate-D-mannose 4,6-deydratase, GMD, and guanosine 5'-diphosphate-Lfucose synthetase GFS.

53. (currently amended) The cell of claim 39 comprising genes encoding galactokinase, GalK, galactose-1-phosphate uridylyltransferase, GalT, and glucose-1-phosphate uridylyltransferase GalU.

- 54. (currently amended) The cell of claim 39 comprising a gene encoding <u>nucleotide</u> diphosphate kinase Ndk.
- 55. (currently amended) The cell of claim 53 comprising a gene encoding polyphosphate kinase PpK,
- 56. (currently amended) The cell of claim 53 comprising a gene encoding <u>pyruvate</u> kinase <u>PykF</u>.
- 57. (currently amended) The cell of claim 53 comprising genes encoding <u>pyruvate</u> oxidase PoxB, nucleotide diphosphate kinase Ndk, and <u>pyrophosphatase</u> Ppa.
- 58. (currently amended) The cell of claim 39 comprising a gene encoding sucrose synthetase SusA.
- 59. (currently amended) The cell of claim 58 further comprising a gene encoding uridine 5'-diphosphate-galactose 4-epimease or uridine 5'-diphosphate-glucose 4-epimerase GalE.
- 60. (currently amended) The cell of claim 58 further comprising a gene encoding glucosyltransferase GluT.
- 61. (currently amended) The cell of claim 58 further comprising genes encoding uridine 5'-diphosphoglucouronic acid 6-dehydrogenase Ugd and a glucuronyltransferase UGT2B7.
- 62. (previously presented) The cell of claim 39, wherein the one or more glycosyltransferase(s) is selected from the group consisting of a galactosyltransferase, a glucosyltransferase, a N-acetylglucosaminyl transferase, an N-acetylgalactosaminyl transferase, a glucuronyltransferase, a sialyltransferase, a mannosyltransferase, and a fucosyltransferase.

63. (currently amended) The cell of claim 62 wherein the galactosyltransferase is selected from the group consisting of α1,3 galactosyltransferase, β1,4 galactosyltransferase, and α1,4 galactosyltransferase LgtB and LgtC.

- 64. (currently amended) The cell of claim 62, wherein the <u>glycosyltransferase</u> is a glucosyltransferase is selected from the group consisting of LgtF, Alg5, and DUCT.
- 65. (currently amended) The cell of claim 62, wherein the <u>glycosyltransferase</u> is a N-acetylglucosaminyl transferase is <u>LgtA</u>.
- 66. (currently amended) The cell of claim 62, wherein the N-acetylgalactosaminyl transferase is <u>uridine 5'-diphosphate-N-acetylgalactosamine:2'-fucosylgalactoside-α-3-N-acetylgalactosaminyl transferase UDP GalNAe:2' fucosylgalactoside a 3 N-acetylgalactosaminyl transferase.</u>
- 67. (currently amended) The cell of claim 62, wherein the <u>glycosyltransferase</u> is a glucuronyltransferase is UGT2B7.
- 68. (currently amended) The cell of claim 62, wherein the <u>glycosyltransferase</u> is a sialyltransferase is SiaT 0160.
- 69. (currently amended) The cell of claim 62, wherein the <u>glycosyltransferase</u> is a mannosyltransferase is selected from the group consisting of Alg1 and Alg2.
- 70. (currently amended) The cell of claim 62, wherein the fucosyltransferase is selected from the group consisting of α 1,3- fucosyltransferase, α 1,2- fucosyltransferase, and α 1,3/4- fucosyltransferase α 1,3-FueT, α 1,2-FueT, and α 1,3/4-FueT.
- 71. (withdrawn) A method of producing a glycoconjugate of interest in the absence of an exogenously supplied nucleotide triphosphate, the method comprising:

contacting a cell comprising heterologous genes encoding susA, galE, and a glycosyltransferase with a bioenergetic.

72. (withdrawn) The method according to claim 71, wherein the bioenergetic comprises fructose generated within the cell by susA acting on sucrose.

- 73. (withdrawn) The method according to claim 72 further comprising supplying the sucrose to the cell.
- 74. (withdrawn) The method according to claim 71, wherein the glycosyltransferase is a galactosyltransferase.
- 75. (withdrawn) The method according to claim 74, wherein the galactosyltransferase is $\alpha 1,3$ GaIT.
- 76. (withdrawn) The method according to claim 71, wherein the glycosyltransferase is LgtC.